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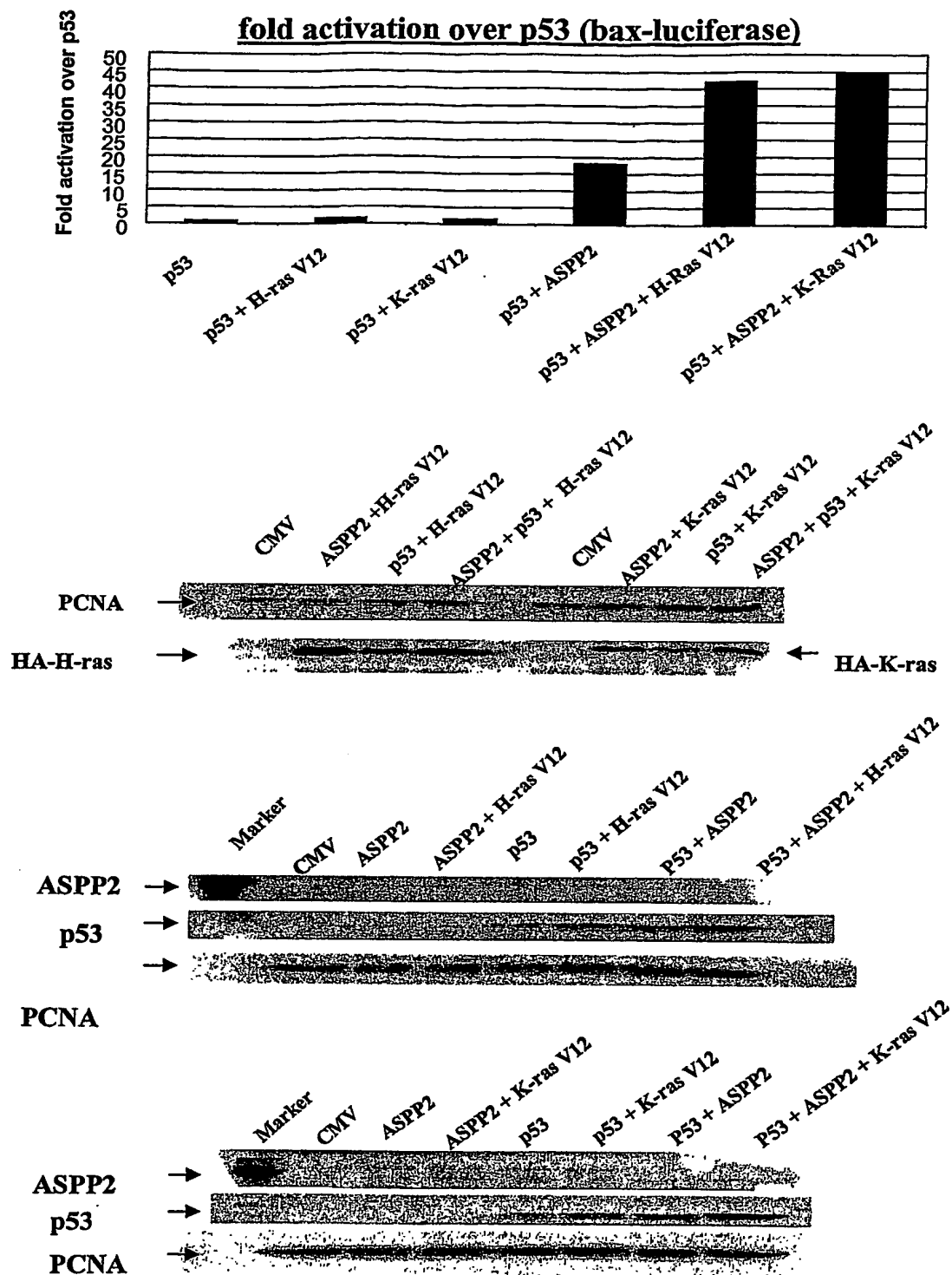
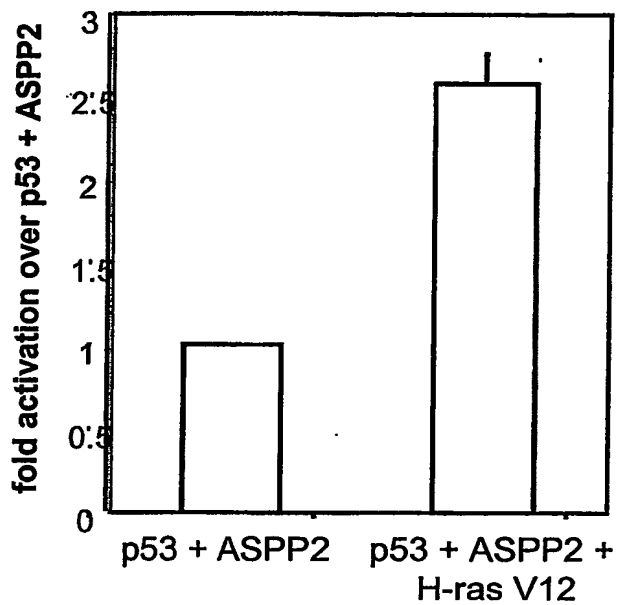


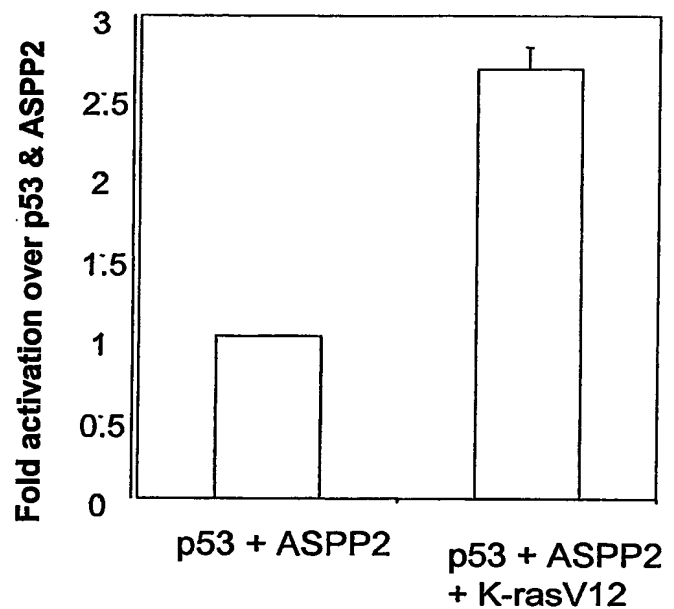
Figure 2

H-ras and K-ras activate ASPP equally

**H-rasV12 activation**  
**of p53 & ASPP2 synergy**



**K-rasV12 activation**  
**of p53 & ASPP2 synergy**



**Figure 3**

Figure 4A

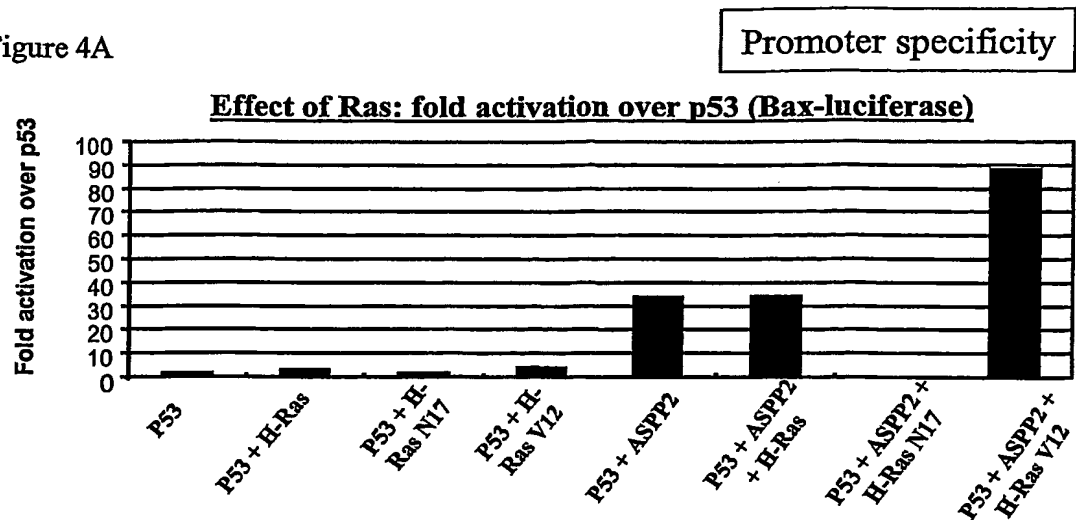


Figure 4B

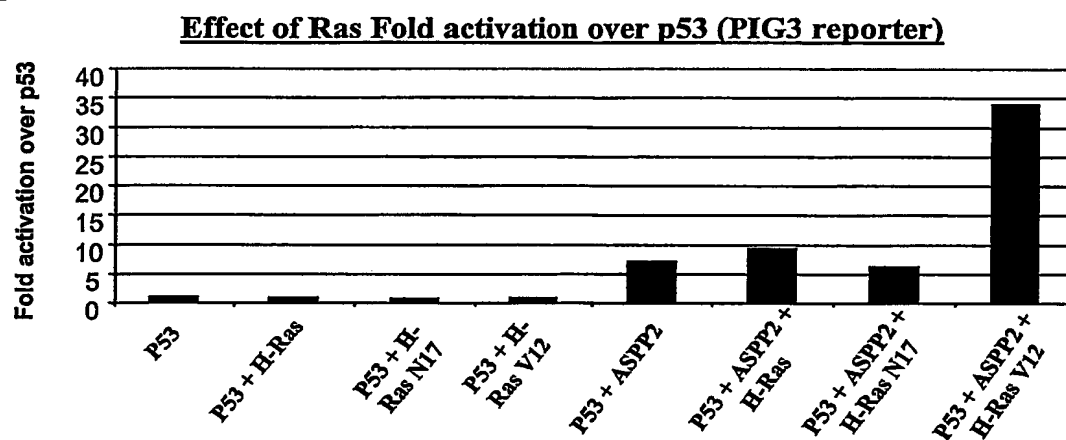


Figure 4C

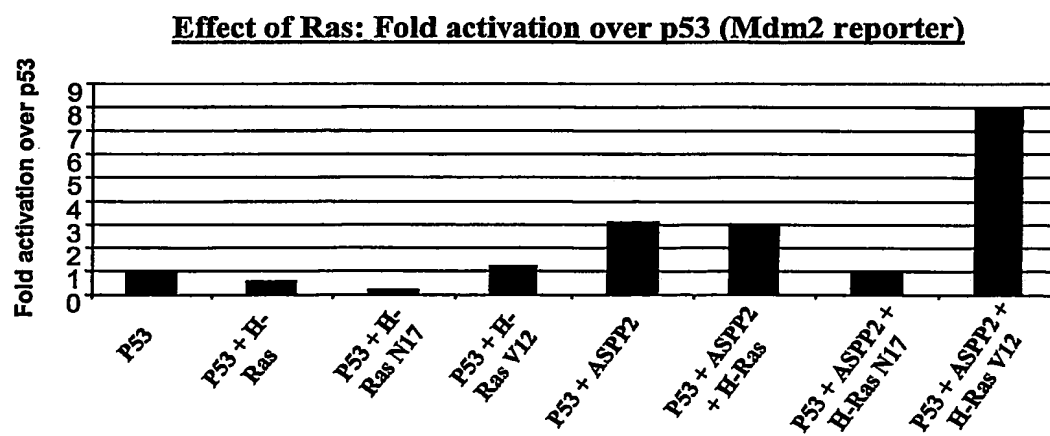


Figure 4D

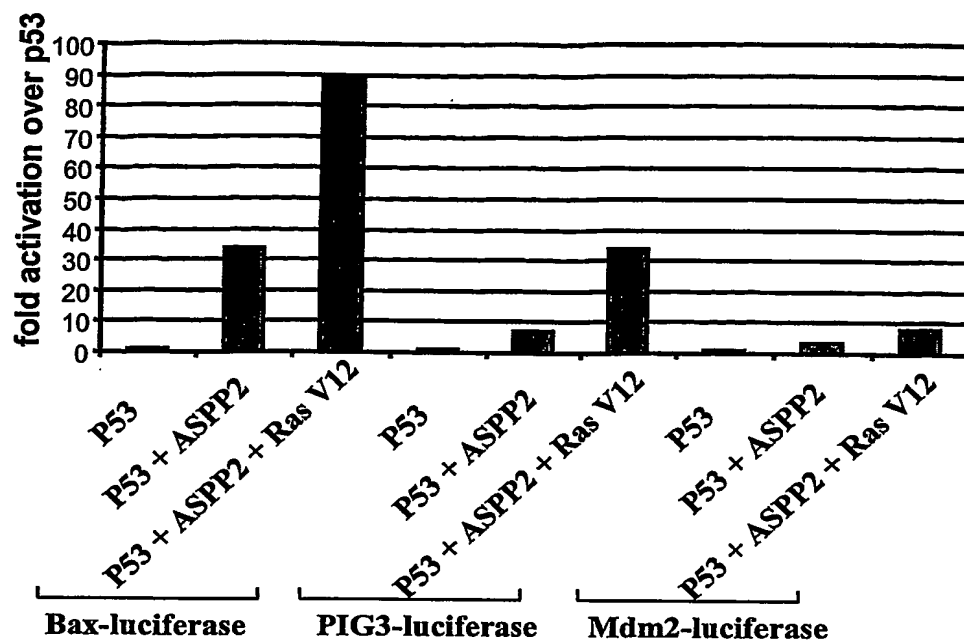
**Effect of rasV12 on transactivation: comparing three reporters**

Figure 4E

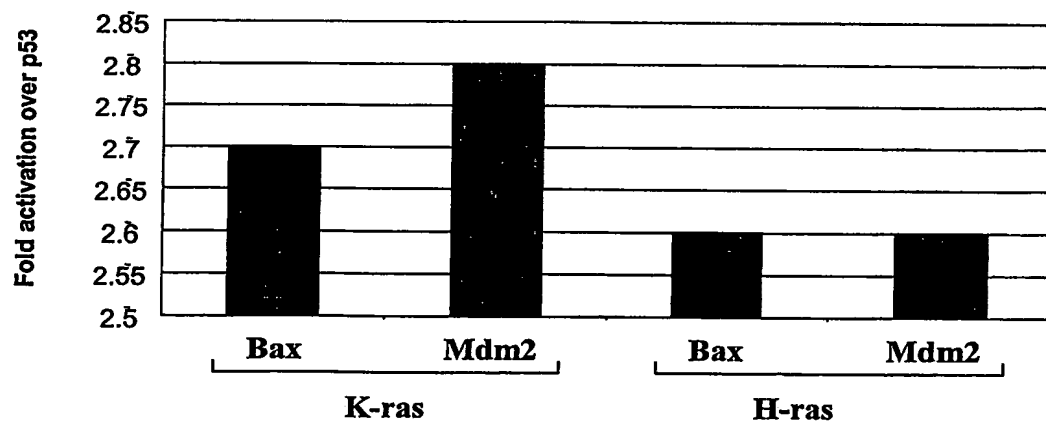
**Promoter specificity of rasV12**

Figure 5A

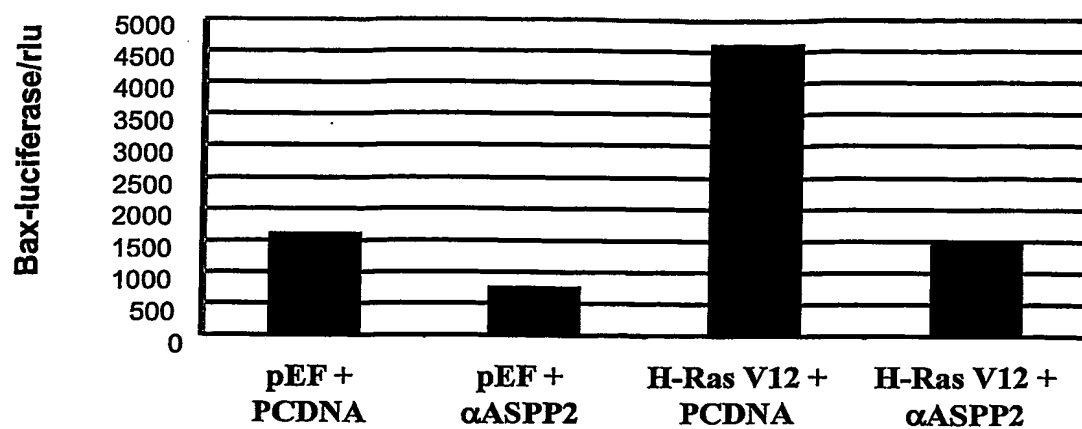
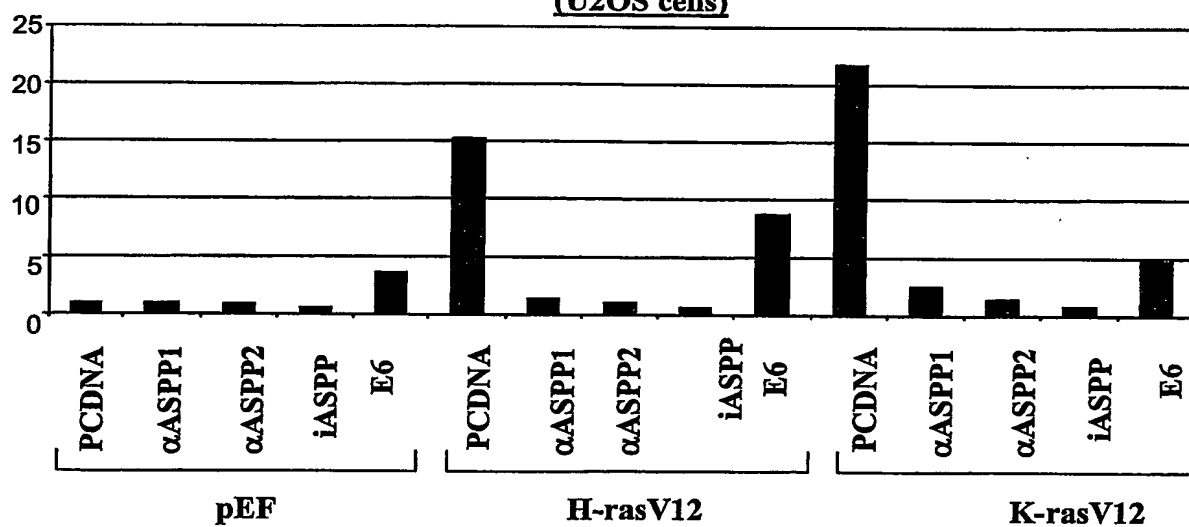
**H-rasV12 activates endogenous ASPP2 to transactivate bax reporter**  
**(U2OS cells)**

Figure 5B

**H- and K-rasV12 activate endogenous ASPP1, ASPP2 and p53 to transactivate bax-reporter**  
**(U2OS cells)**

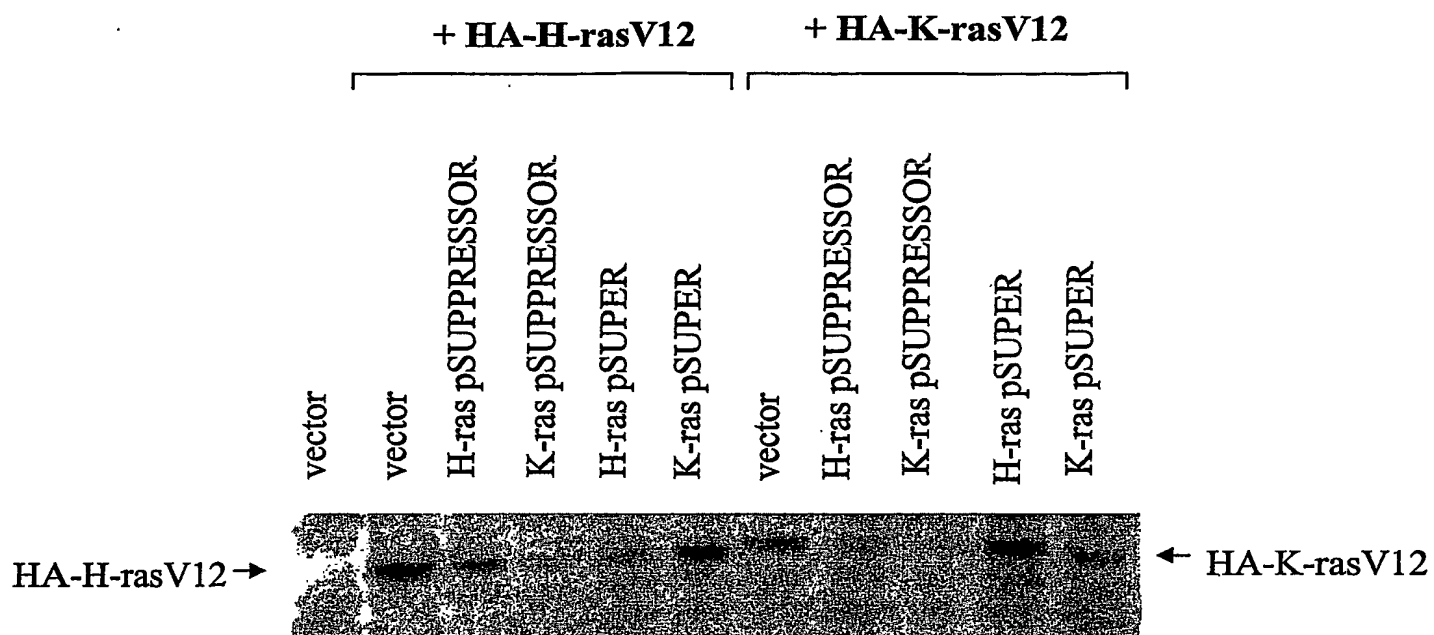


Figure 6

Figure 6B

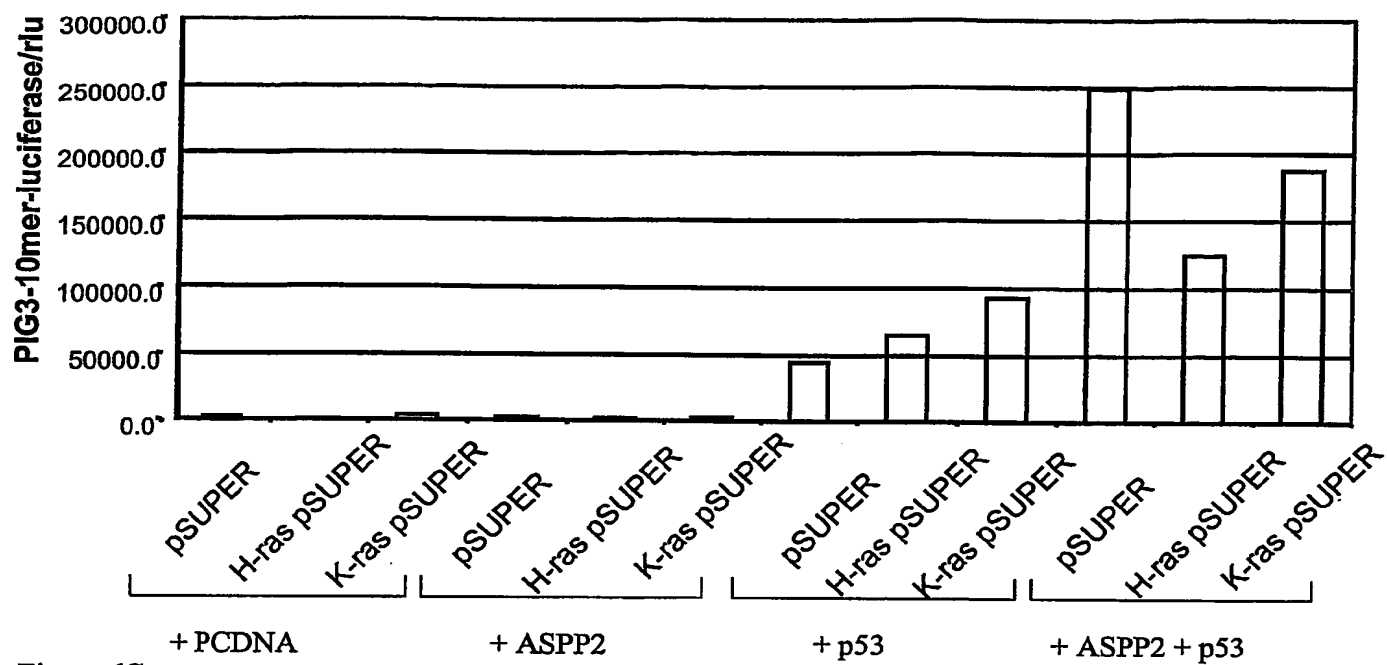


Figure 6C

ASPP2

p53

PCNA

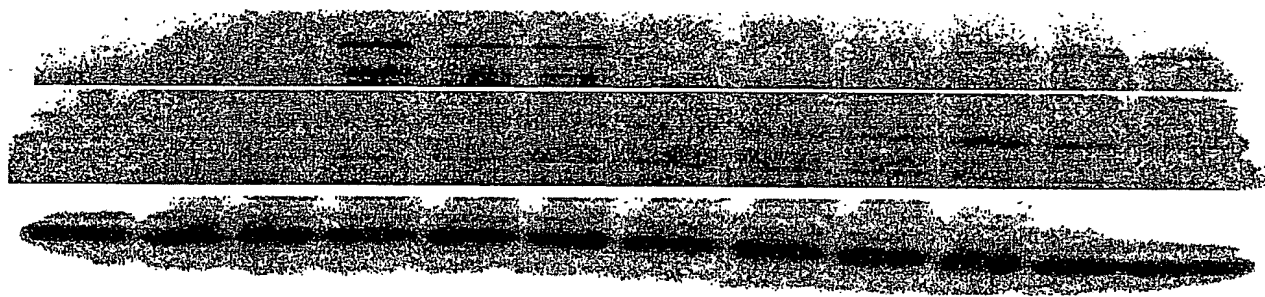




Figure 7B

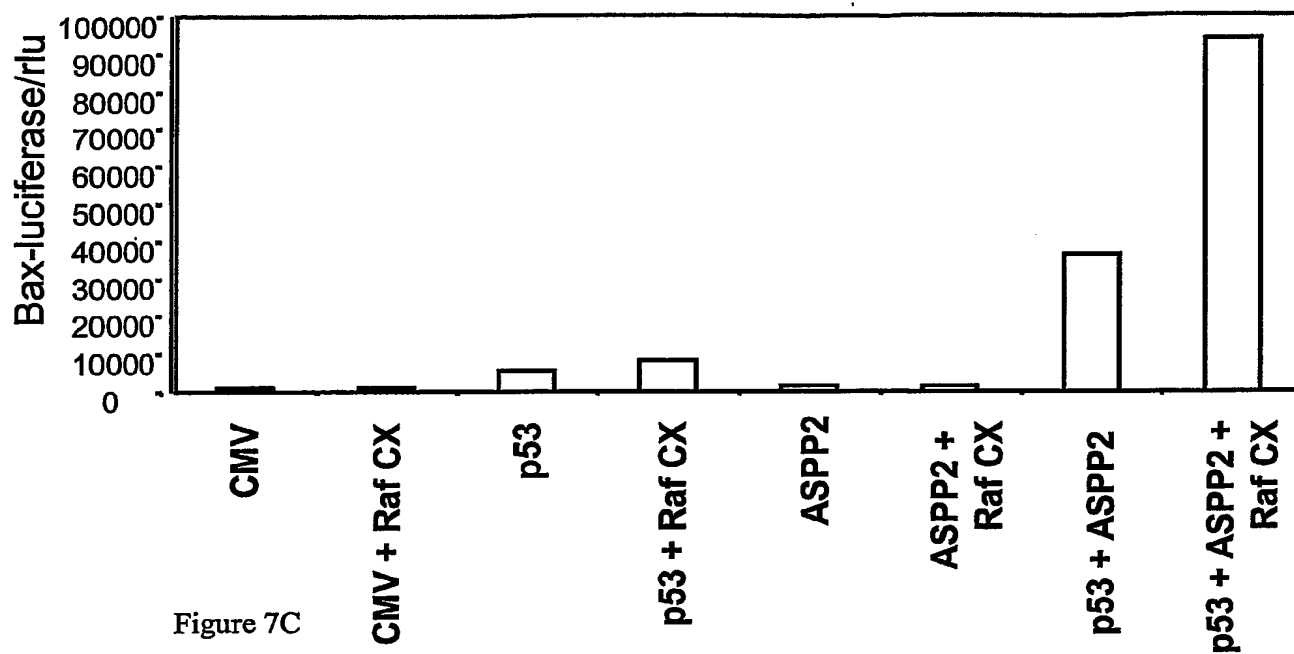


Figure 7C

ASPP2



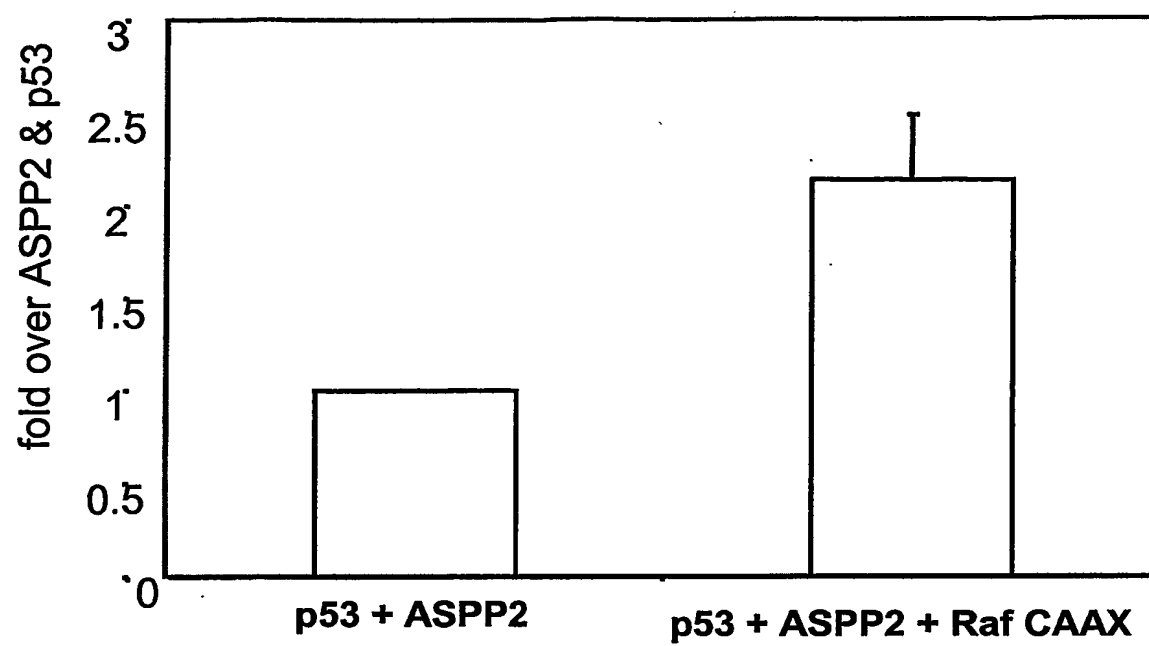
Raf



p53



Figure 7d

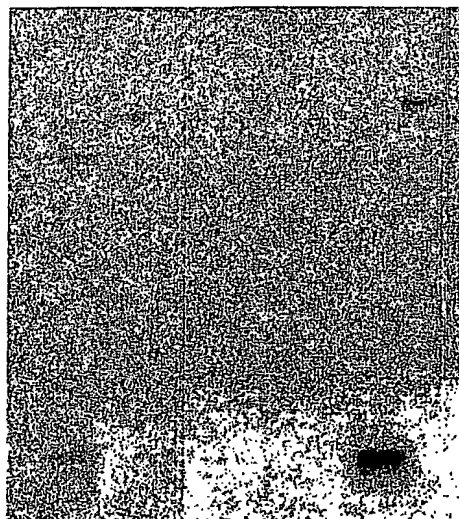


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Figure 8A

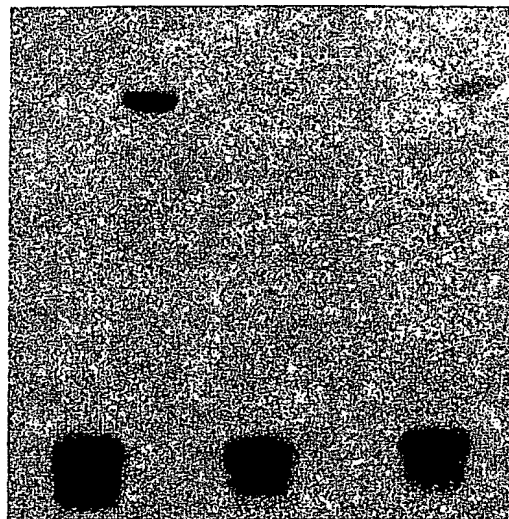
MAPK1 p70S6K p90rsk

H2B ASPP2 H2B ASPP2 H2B ASPP2



PKA PKB p38SAPK

H2B ASPP2 H2B ASPP2 H2B ASPP2



ASPP2

H2B

Figure 8B

PKA p38 SAPK MAPK1 p90rsk

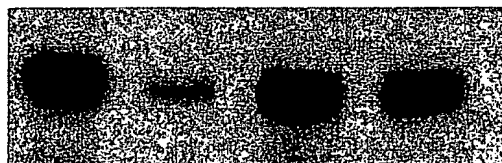


Figure 8C

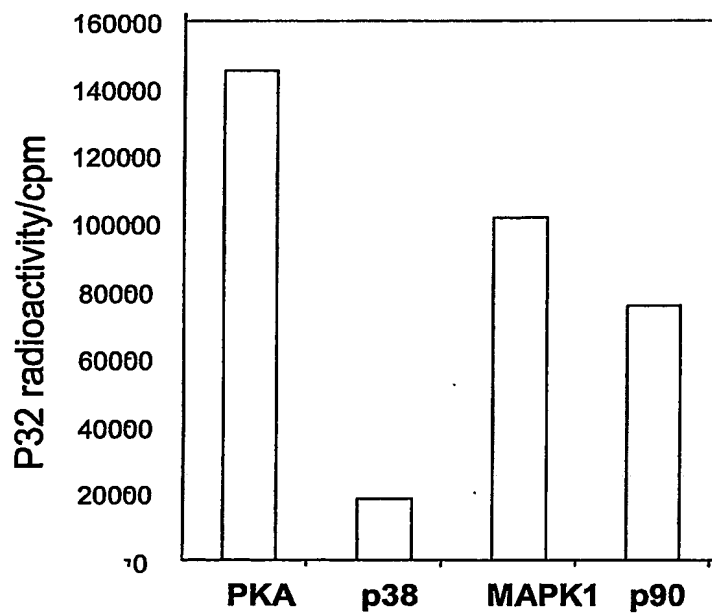


Figure 8D

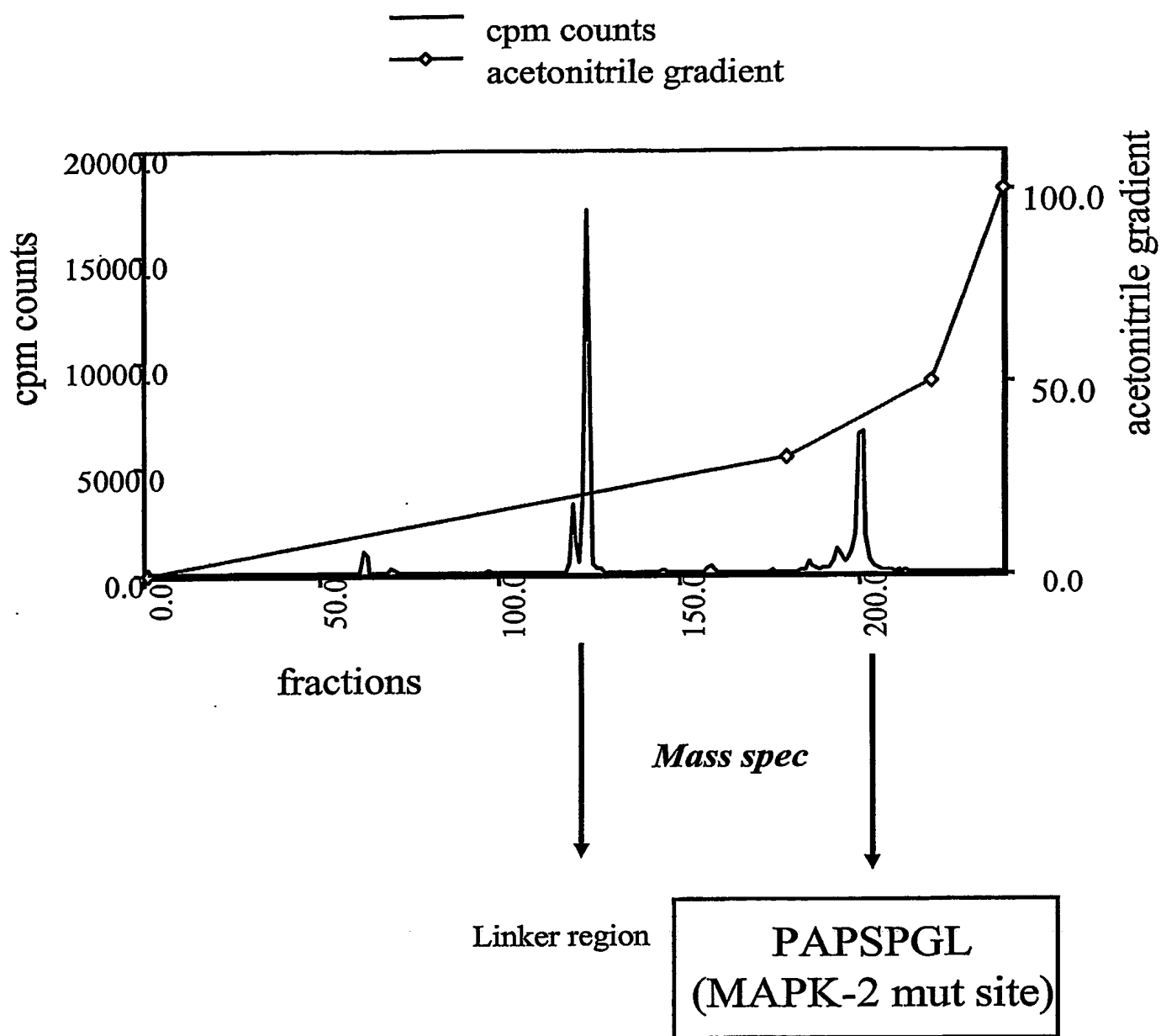


Figure 9

**C-term of ASPP2:**

550 - QPRVLLSPSIPSVGQDQTLSPGSKQESPPAAAVRPFTPQPS  
KDTLLPPFRKPQTVAAASSIYSMYTQQQAPGKNFQQA VQS  
ALTKTHTRGPHFSSVYGKPVIAAAQNQQHPENIYSNSQ  
GKPGSPEPETEPVSSVQENHENERIPRPLSPTKLLPFLSNP  
YRNQSDADLEALRKKLSNAPRPLKKRSSITEPEGPNGPNI  
QKLLYQRTTIAAMETSVPSYPSKSASVTASSESPVEIQNP  
YLHVEPEKEVVSLVPESLSPEDVGNASTENS DMPAPSPGL  
DYEPEGVPDNSPNLQNNPE - 849

**S** — MAPK sites  
**SS** — PKA site

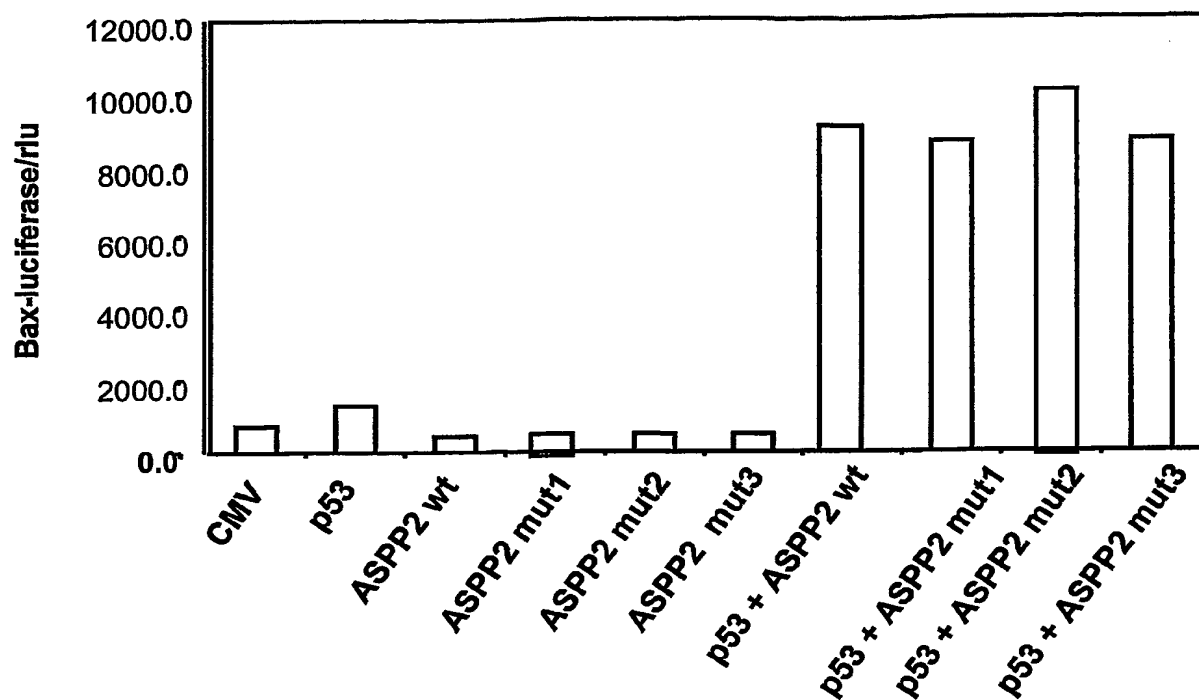
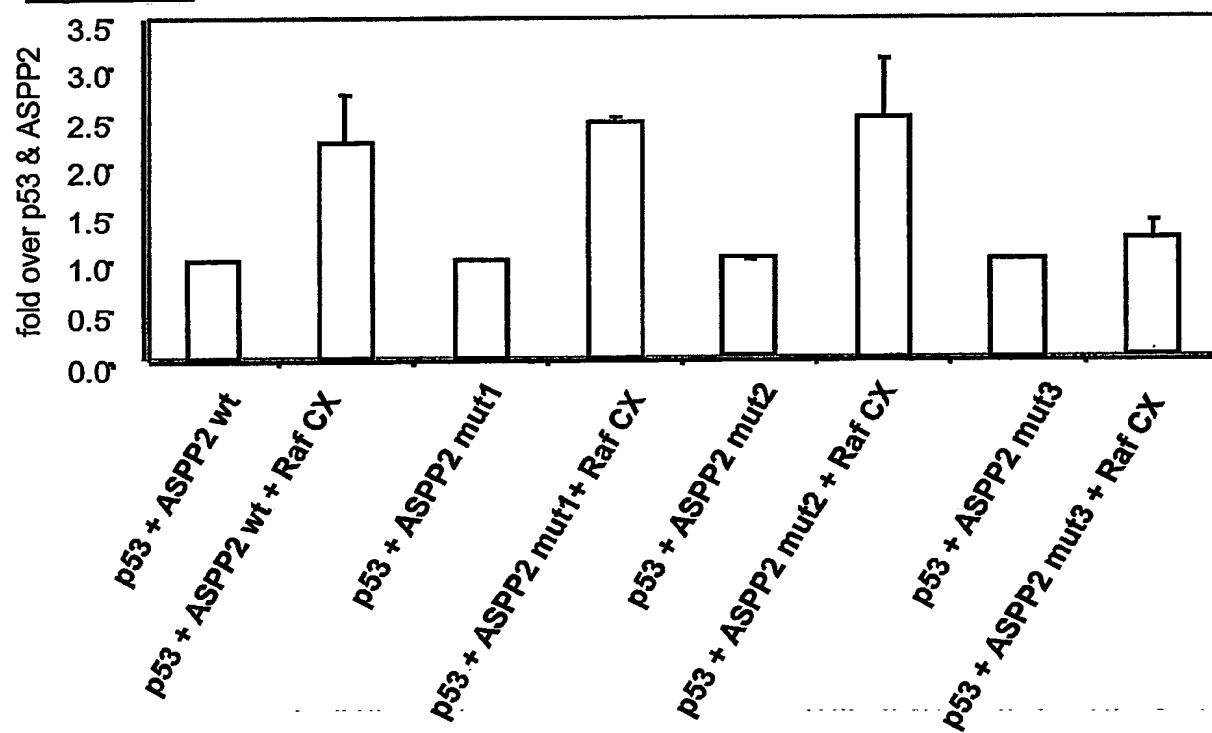
**Figure 10A****Figure 10B**

Figure 11A

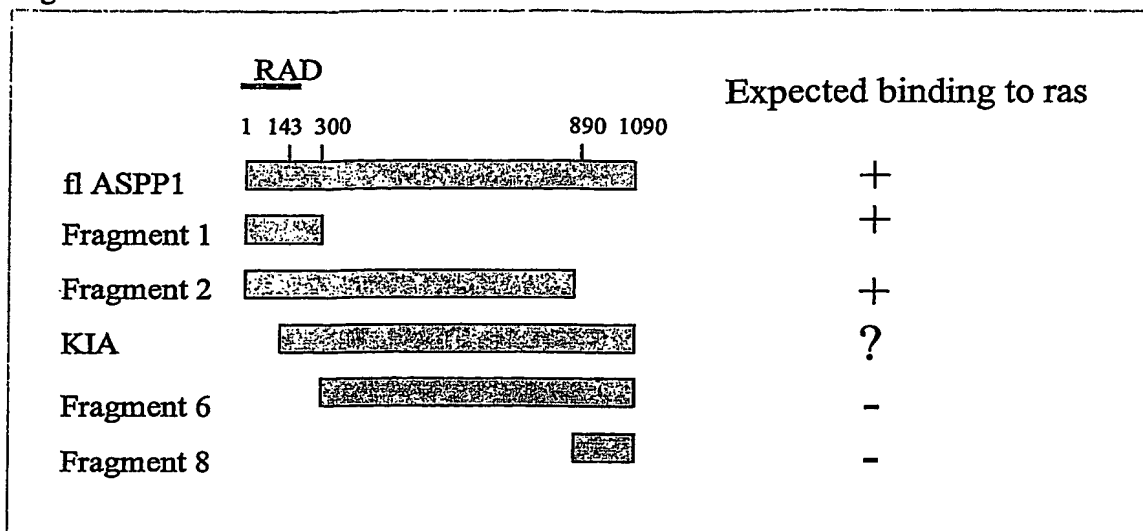


Figure 11B

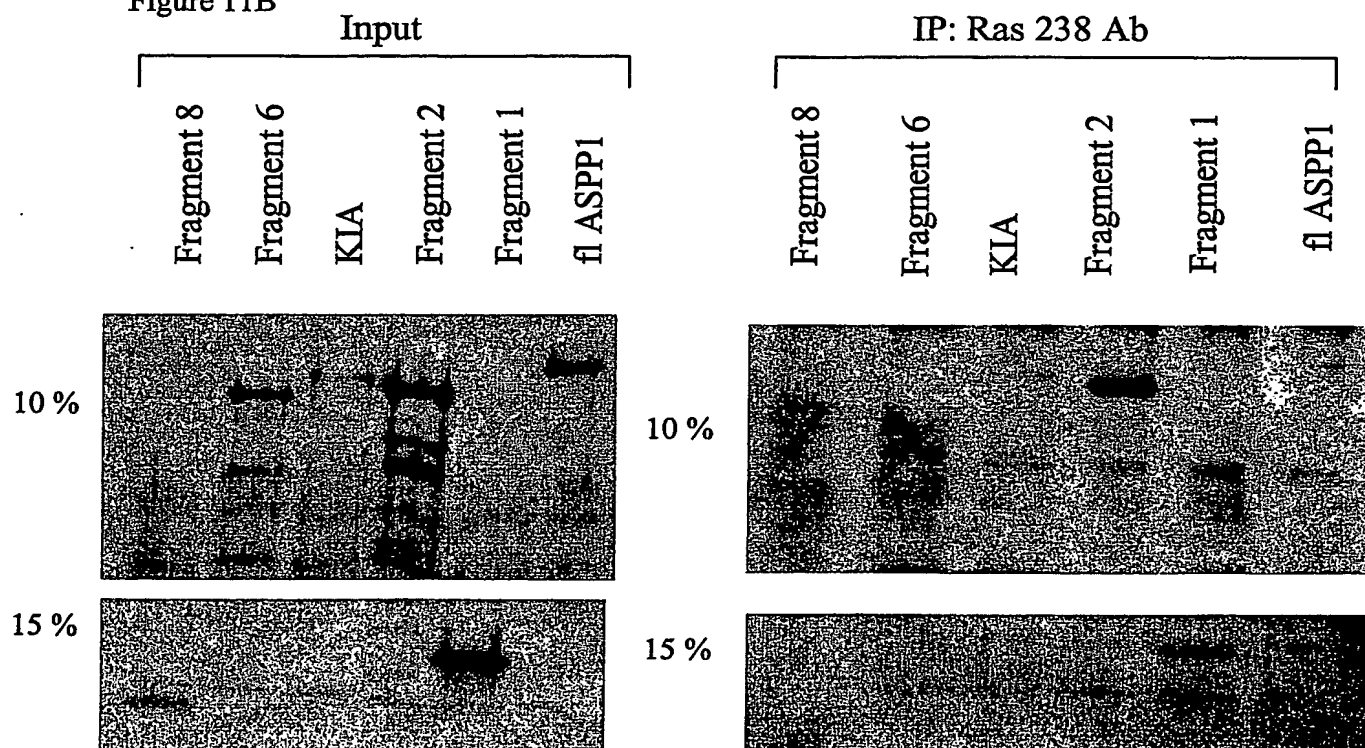


Figure 12

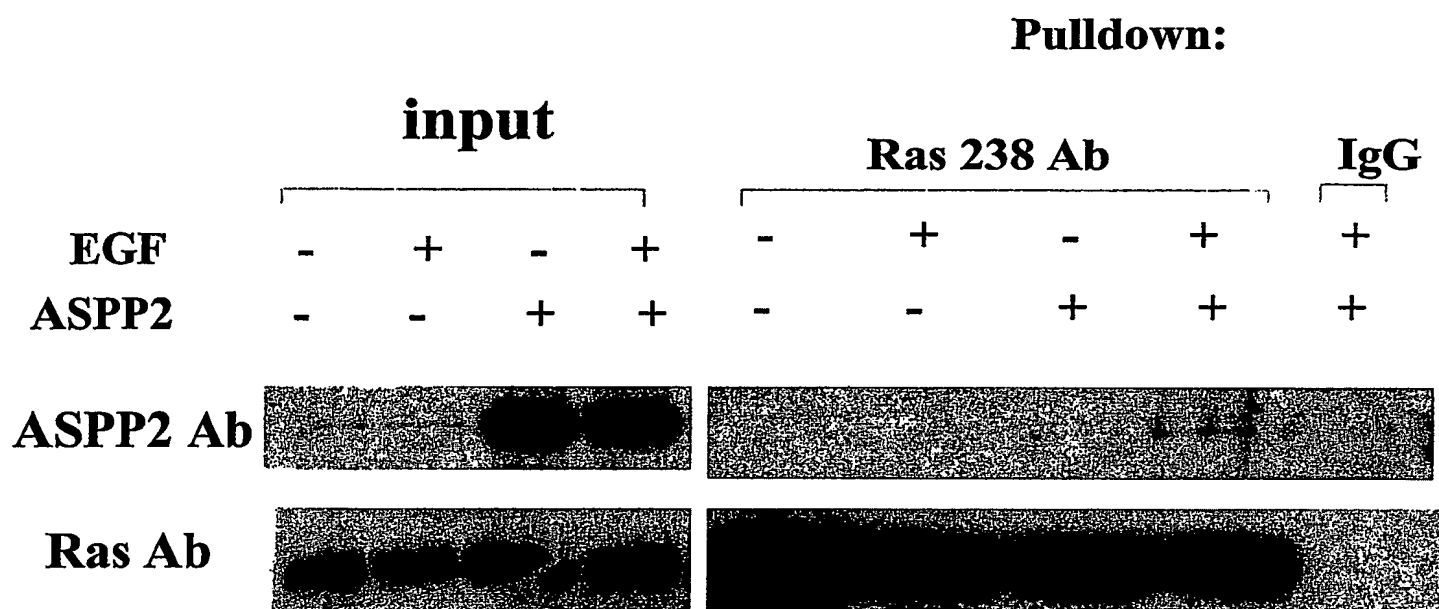
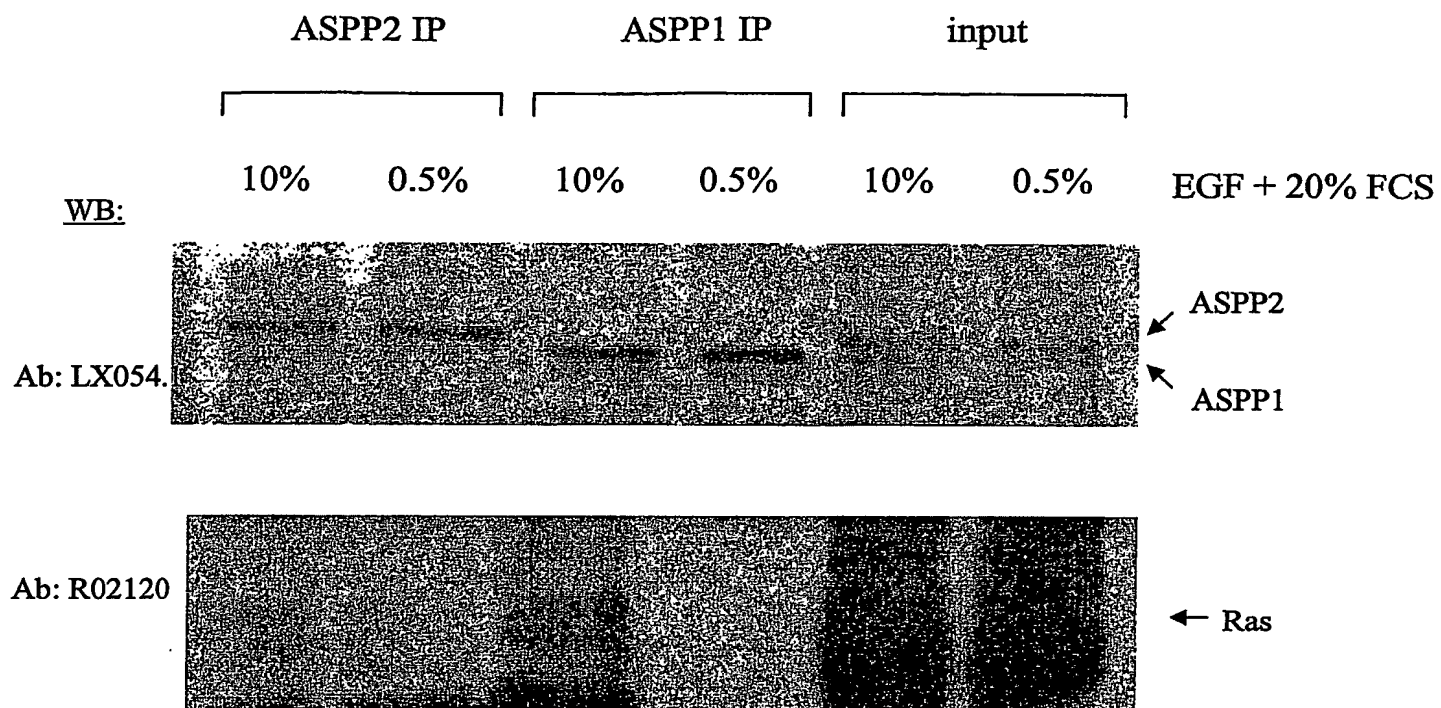
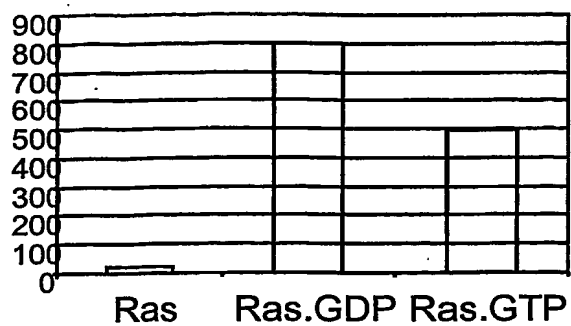


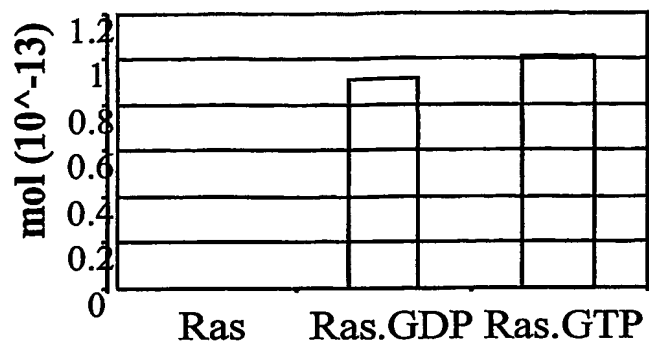
Figure 13



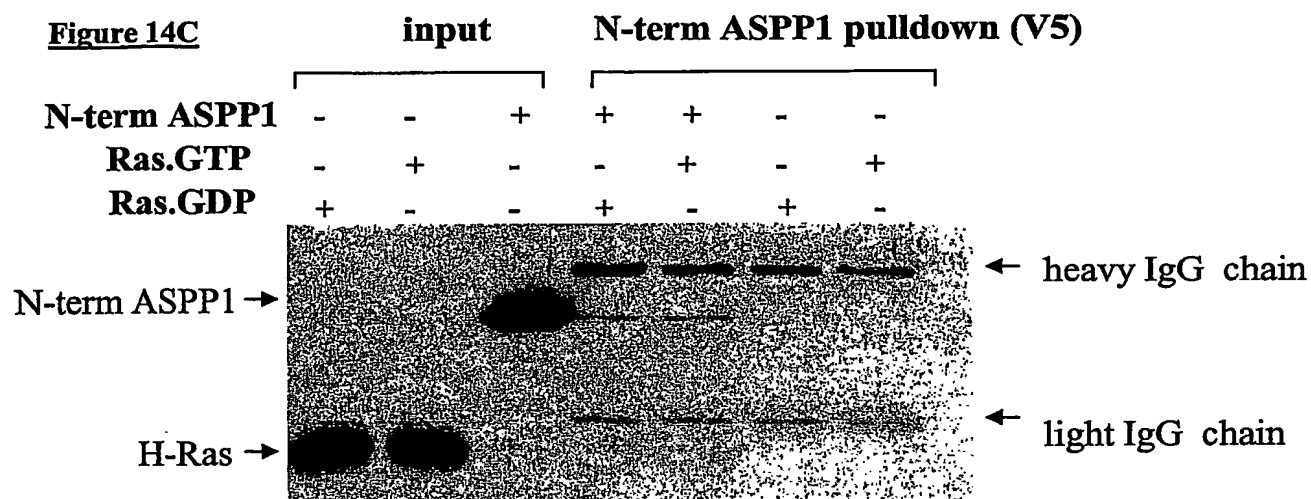
**Figure 14A**



**Figure 14B**



**Figure 14C**



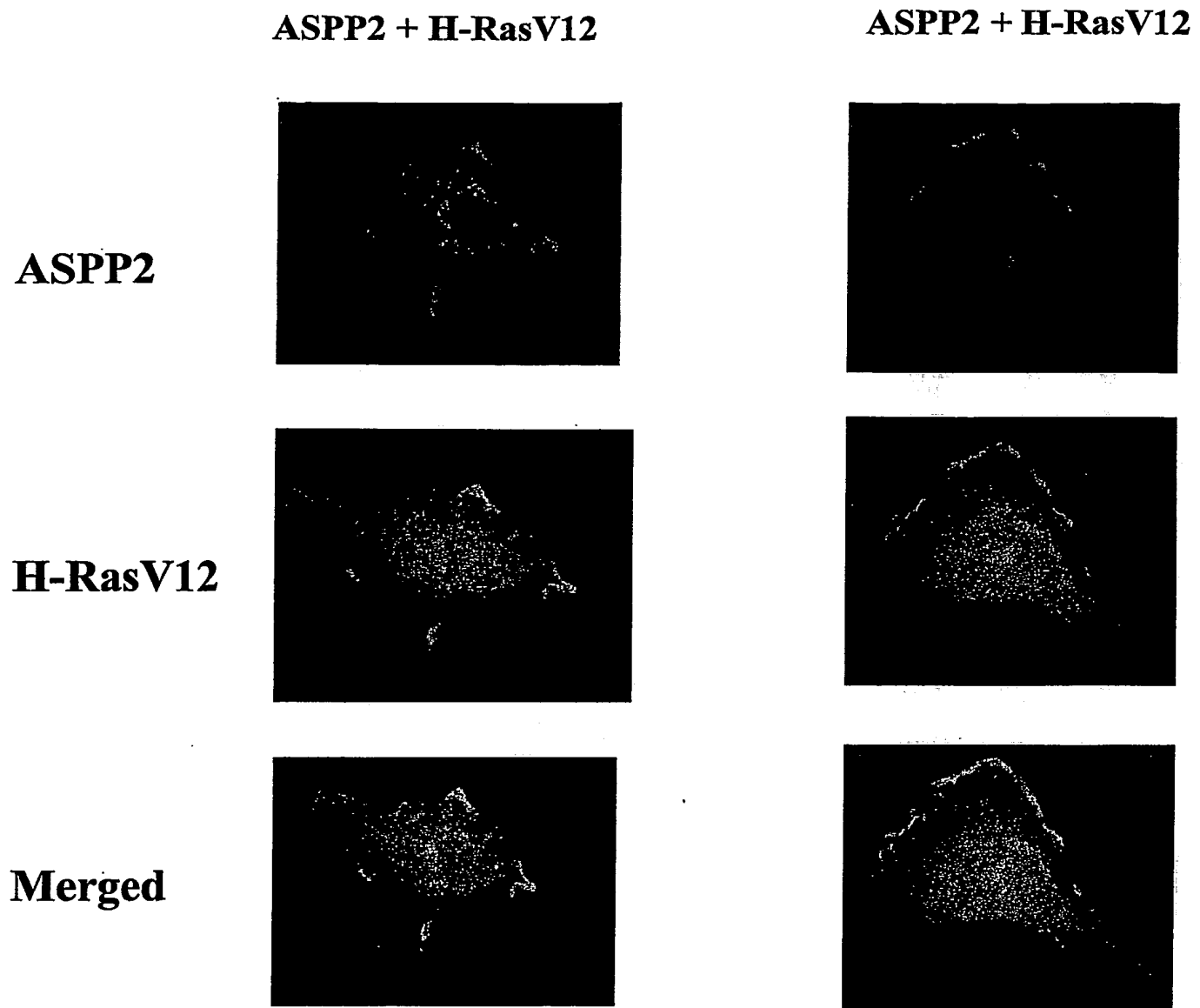


Figure 15

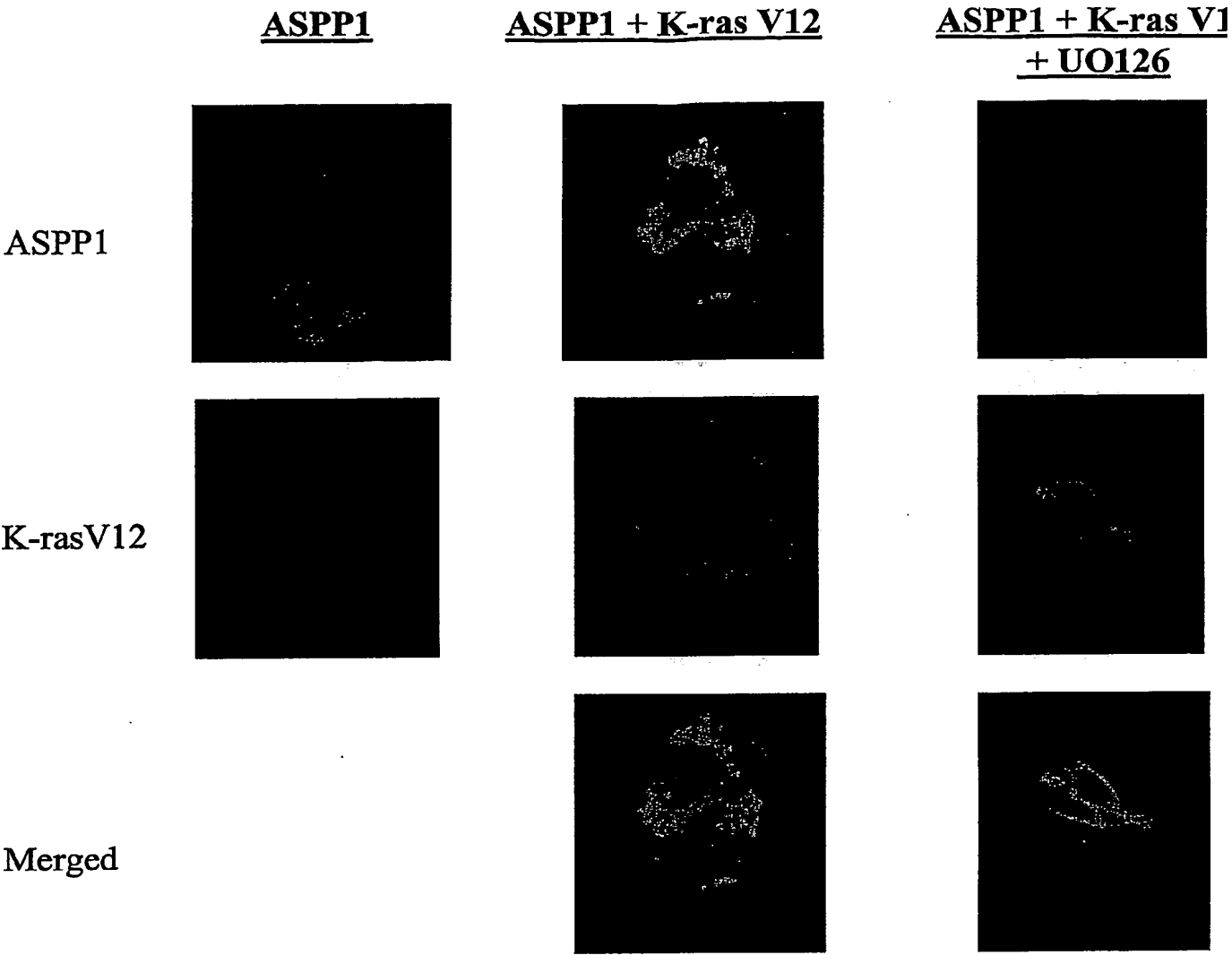


Figure 16

Figure 17a

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Figure 17b

GTCACGAGCGTCGAAGAGACAAAGCCGCGTCAGGGGGCCCGCCGGGGGAGCCCGGGGCTTGTTGGTGCCCCAGC  
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CT

Figure 17c

MMPMILT VFLSNNEQILTEVPITPETTCRDVVEFCKEPGECSCHLAEBVWRGNERPIPFDHMMYEH LQIWGPRREEVKFFL  
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PSESTEKEPEQDGPAPADGSTVESLPRPLSPTKLTPIVHSPLRYQSDADLEALRRKLANAPRPLKKRSSITEPEGPGGP  
NIQKLLYQRFNTLAGGMEGTPFYQPSPSQDFMGT LADVDNGNTNANGNLEELPPAQPTAPLPAEPAPSSDANDNBLSPE  
PEELICPQTTHQTAEPAEDNNNNVATVPTTEQIPSPVAEAPSPGEEQVPPAPLP PASHPPATSTNKRTNLKKPNSERTGH  
GLRVRFNPLALLLDASLEGEFDLVQRIIYEVEDPSKFNDEGITPLHNAVCAGHHHIVKFLLD FGVNVNAADSDGWTP LHC  
AASCNSVHLCKQLVESGA AIFASTISDIETAADKCEMEEGYIQCSQFLYGVQEKLGVMNKGVA YALWDYEAQNSDELSF  
HEGDALTILRRKDESETEWWARLG DREGYVPKNLLGLYPRIKPRQRTLA

Figure 17d

MMPMFLTVYLSNNEQHFTVEPVTPETICRDVVDLCKEPGESDCHLAEVWCGSERPVADNERMFDVLQRFQSQRNEVRFFL  
RHERPPGRDIVSGPPRSQDPSLKRNGVKVPGEYRRKENGVNSPRMDLTLAELQEMASRQQQIEAQQQLLATKEQRLKFLK  
QQDQRQQQVAEQEKLRLEIAENQEAQLKKVRALKGHVEQKRLSNGKLVBEIEQMNNLFQKKQRELVLAVSKVEELTR  
QLEMLKNGRIDSHHDNQSAVAELDRLYKELQLRNKLNQEQNAKLQQRECLNKRNSEVAVMDKRVNELRDRLWKKKAALQ  
QKENLPVSSDGNLPQQAASAPSRVAAVGPYIQSSTMPRMPSPRPELLVKPALPDGSLVIOASEGPMKIQTLPNMRSGAASQ  
TKGSKIHPVGPDWSPSNADLFPSQGSASVPQSTGNALDQVDDGEVPLREKEKKVRPFMSMFDVAVDQSNAPPSPFGTLRKNQS  
SEDILRDAQVANKNVAKVPPVPTPKQINLPYFGQTNQPPSDIKPDGSSQQLSTVVPSMGTKPKPAGQQPRVLLSPSIP  
SVGQDQTLSPGSKQESPPAAVRPFTFPQPSKDTLLPFFRKPTVAASSIYSMTQQQAPGKNFQQAVQSALTTKTHTRGPH  
FSSVYGKPVIAAAQNNQHPENIYNSQGKPGSPETEPVSSVQENHENERIPRPLSPTKLLPFLSNFYRNQSDADLEA  
LRKKLSNAPRPLKKRSSITEPEGPNPNIQKLLYQRTTIAAMETISVPSYPSKSASVTASSESPVEIQNPYLHVEPEKEV  
VSLVPESLSPEDVGNASTENSMDPAPSPGLDYEPEGVDPNSPNLQNNPBEENPEAPHVLDVYLEEYPYPYPYPYPSPGEPE  
GPGEDSVSMRPPFITGQVSLPPGKRTNLRKTGSERIAHGMRVKFNPLALLLDSSLEGEFDLVQRIIYEVDPSLPNDEGI  
TALHNAVCAHTEIVKFLVQFGVNVNAADSDGWTPLHCAASCNNVQVCKFLVESGAAVFAMTYSDMQTAADKCEEMEegy  
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KPRQRSIA

Figure 18a

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GTGCCTGTTGGACATCCTGGATAACGCCGGCCAGGAGGAGTACAGCGCCA  
TGCGGGACCAGTACATGCGCACCGGGGAGGGCTTCCTGTGTGTGTTTGCC  
ATCAACAACACCAAGTCTTTTGAGGACATCCACCAGTACAGGGAGCAGAT  
CAAACGGGTGAAGGACTCGGATGACGTGCCCATGGTGCTGGTGGGGAAC  
AAGTGTGACCTGGCTGCACGCACTGTGGAATCTCGGCAGGCTCAGGACCT  
CGCCCGAAGCTACGGCATCCCCTACATCGAGACCTCGGCCAAGACCCGGC  
AGGGAGTGGAGGATGCCTTCTACACGTTGGTGCGTGAGATCCGGCAGCAC  
AAGCTGCGGAAGCTGAACCCTCCTGATGAGAGTGGCCCCGGCTGCATGAG  
CTGCAAGTGTGTGCTCTCCTGA

Figure 18b

MTEYKLVVVGAGGVGKSALTIQLIQNHFVDEYDPTIEDSYRKQVVIDGETCL  
LDILD TAGQEEYSAMRDQYMRTGEGFLCVFAINNTKSFEDIHQYREQIKRVK  
DSDDVPMVLVGNKCDLAARTVESRQAQDLARSYGIPYIETSAKTRQGVEDAF  
YTLVREIRQHKLRKLNPPDESGPGCMSCKCVLS

Figure 18c

ATGACGGAATATAAGCTGGTGGTGGTGGGCGCCGTCGGTGTGGGCAAGA  
GTGCGCTGACCATCCAGCTGATCCAGAACCATTTTGTGGACGAATACGAC  
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GTGCCTGTTGGACATCCTGGATACCGCCGGCCAGGAGGAGTACAGCGCCA  
TGCGGGACCAGTACATGCGCACCGGGGAGGGCTTCCTGTGTGTGTTTGCC  
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CAAACGGGTGAAGGACTCGGATGACGTGCCCATGGTGCTGGTGGGGAAC  
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CGCCCGAAGCTACGGCATCCCCTACATCGAGACCTCGGCCAAGACCCGGC  
AGGGAGTGGAGGATGCCTTCTACACGTTGGTGCGTGAGATCCGGCAGCAC  
AAGCTGCGGAAGCTGAACCCTCCTGATGAGAGTGGCCCCGGCTGCATGAG  
CTGCAAGTGTGTGCTCTCCTGA

Figure 18d

MTEYKLVVVGAVGVGKSALTIQLIQNHFVDEYDPTIEDSYRKQVVIDGETCL  
LDILDTAGQEEYSAMRDQYMRTGEGFLCVFAINNTKSFEDIHQYREQIKRVK  
DSDDVPMVLVGNKCDLAARTVESRQAQDLARSYGIPYIETSAKTRQGVEDAF  
YTLVREIRQHKLRKLNPPDESGPGCMSCKCVLS-

Figure 18e

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AACAAATAGAGGATTCCTACAGGAAGCAAGTAGTAATTGATGGAGAAACC  
TGTCTCTTGGATATTCTCGACACAGCAGGTCAAGAGGAGTACAGTGCAAT  
GAGGGACCAGTACATGAGGACTGGGGAGGGCTTTCTTTGTGTATTTGCCA  
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ATGTGATTTGCCTTCTAGAACAGTAGACACAAAACAGGCTCAGGACTTAG  
CAAGAAGTTATGGAATTCCTTTTATTGAAACATCAGCAAAGACAAGACAG  
GGTGTGATGATGCCTTCTATACATTAGTTTCGAGAAATTCGAAAACATAA  
AGAAAAGATGAGCAAAGATGGTAAAAAGAAGAAAAAGAAGTCAAAGAC  
AAAGTGTGTAATTATGTAA

Figure 18f

MTEYKLVVVGAGGVGKSALTIQLIQNHFVDEYDPTIEDSYRKQVVIDGETCL  
LDILDTAGQEEYSAMRDQYMRTGEGFLCVFAINNTKSFEDIHHYREQIKRVK  
DSEDVPMVLVGNKCDLPSRTVDTKQAQDLARSYGIPFIETSAKTRQGVDDAF  
YTLVREIRKHKEKMSKDGKKKKKKSKTKCVIM-

Figure 18g

ATGACTGAATATAAACTTGTGGTAGTTGGAGCTGTCGGCGTAGGCAAGAG  
TGCCTTGACGATACAGCTAATTCAGAATCATTTTGTGGACGAATATGATCC  
AACAATAGAGGATTCTACAGGAAGCAAGTAGTAATTGATGGAGAAACC  
TGTCTCTTGGATATTCTCGACACAGCAGGTCAAGAGGAGTACAGTGCAAT  
GAGGGACCAGTACATGAGGACTGGGGAGGGCTTTCTTTGTGTATTTGCCA  
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AAAAGAGTTAAGGACTCTGAAGATGTACCTATGGTCCTAGTAGGAAATAA  
ATGTGATTTGCCTTCTAGAACAGTAGACACAAAACAGGCTCAGGACTTAG  
CAAGAAGTTATGGAATTCCTTTTATTGAAACATCAGCAAAGACAAGACAG  
GGTGTTGATGATGCCTTCTATACATTAGTTTCGAGAAATTCGAAAACATAA  
AGAAAAGATGAGCAAAGATGGTAAAAAGAAGAAAAAGAAGTCAAAGAC  
AAAGTGTGTAATTATGTAA

Figure 18h

MTEYKLVVVGAVGVGKSALTIQLIQNHFVDEYDPTIEDSYRKQVVIDGETCL  
LDILDTAGQEEYSAMRDQYMRTGEGFLCVFAINNTKSFEDIHHYREQIKRVK  
DSEDVPMVLVGNKCDLPSRTVDTKQAQDLARSYGIPFIETSAKTRQGVDDAF  
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Figure 19a

```
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caccagacct actgccagag aaccctgagg gagataaaaa tcttactgcg cttcagacat
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taa
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Figure 19b

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TDLYKLLKTQHLSNDHICYFLYQILRGLKYIHSANVLHRDLKPSNLLLNTTCDLKI
CDFGLARVADPDHDHTGFLT EYVATR WYRAPEIMLNSKGYTKSIDIWSVGCILA
EMLSNRPIFP GKHYLDQLKHILGILGSPSQEDLNCIINLKARNYLLSLPHKNKVPW
NRLFPNADSKALDLLDKMLTFNPHKRIEVEQALAH PYLEQYYDPSDEPIAEAPFK
FDMELDDL PKEKLKELIFEETARFQPGYRS
```

Figure 20a

```

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2581 cctgtgtgct gcgaaggacg agacttctc ttgaacagtg tgctgttgta aacatatttg
2641 aaaactatta ccaataaagt tttgttttaa aaaaaaaaaa aaaaa

```

Figure 20b

```

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EPHARFYAAQIVLTFEYLSLDLIYRDLKPENLLIDQQGYIQVTDGFGAKRVKGRWTLCGTPEYLAPEI
ILSKGYNKAVDWWALGVLIYEMAAGYPFFADQPIQIYEKIVSGKVRFP SHFSSDLKDLLRNLLQVDLTK
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F

```

Figure 21a

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CATTCTTCTGGAGCTGGAGGCACCCCTCAAGATCTGCGGTGACATACACG  
GCCAGTACTACGACCTTCTGCGACTATTTGAGTATGGCGGTTTCCCTCCCG  
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TTGGAGACCATCTGCCTGCTGCTGGCCTATAAGATCAAGTACCCCGAGAA  
CTTCTTCCTGCTCCGTGGGAACCACGAGTGTGCCAGCATCAACCGCATCTA  
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CCTTCACTGACTGCTTCAACTGCCTGCCCATCGCGGCCATAGTGGACGAA  
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GCAGATTCGGCGGATCATGCGGCCACAGATGTGCCTGACCAGGGCCTGC  
TGTGTGACCTGCTGTGGTCTGACCCTGACAAGGACGTGCAGGGGCTGGGGC  
GAGAACGACCGTGGCGTCTCTTTTACCTTTGGAGCCGAGGTGGTGGCCAA  
GTTCTCCACAAGCAGACTTGGACCTCATCTGCCGAGCACACCAGGTGG  
TAGAAGACGGCTACGAGTTCTTTGCCAAGCGGCAGCTGGTGACACTTTTC  
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TGTGGACGAGACCCTCATGTGCTCTTTCCAGATCCTCAAGCCCGCCGACA  
AGAACAAGGGGAAGTACGGGCAGTTCAGTGGCCTGAACCCTGGAGGCCG  
ACCCATCACCCACCCCGCAATTCCGCCAAAGCCAAGAAATAG

Figure 21b

MSDSEKLNLDSEIIGRLLEVQGSRPQGNVQLTENEIRGLCLKSREIFLSQPILLEL  
EAPLKICGDIHGQYYDLLRLFYGGFPPESENFLFLGDYVDRGKQSLETICLLL  
AYKIKYPENFFLLRGNHECASINRIYGFYDECKRRYNIKLWKTFTDCFNCLPIA  
AIVDEKIFCCHGGLSPDLQSMEQIRRMPTDVPDQGLLCDLLWSDPDKDVQ  
GWGENDRGVSTFGAEVVAKFLHKHDLDLICRAHQVVEDGYEFFAKRQLVT  
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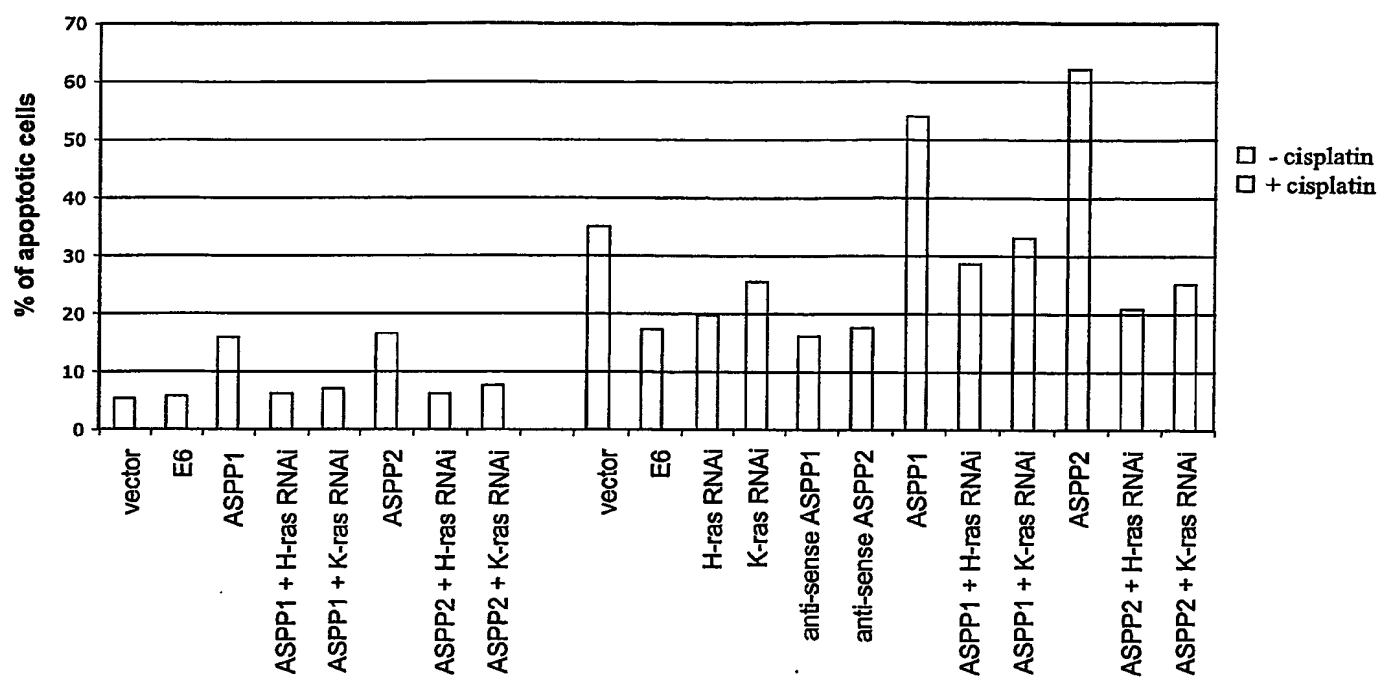
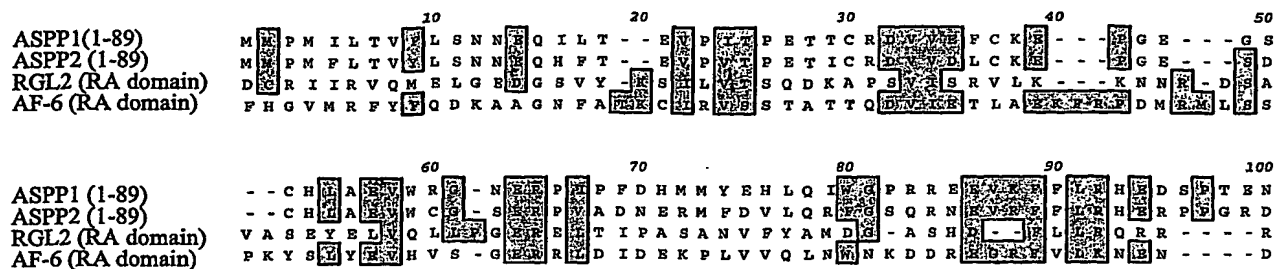
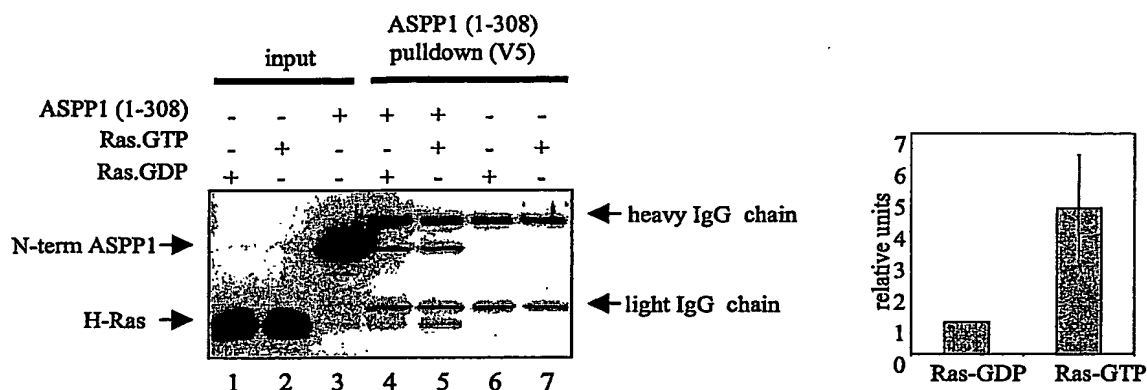


Figure 22

23  
A

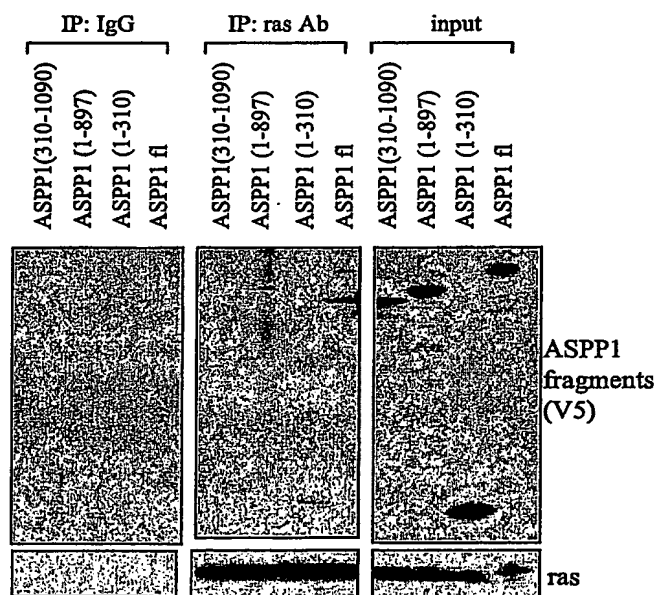


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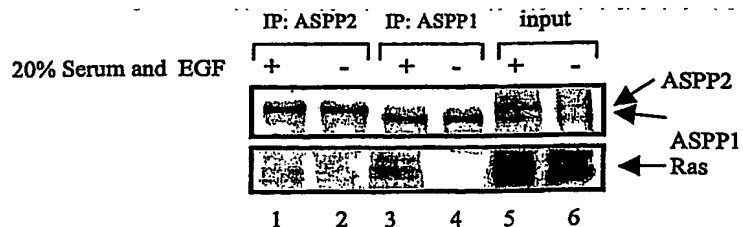


23  
C

	RAD	Contains RAD
ASPP1 fl	1 143 310 897 1090	+
ASPP1 (1-310)		+
ASPP1 (1-897)		+
ASPP1 (310-1090)		-

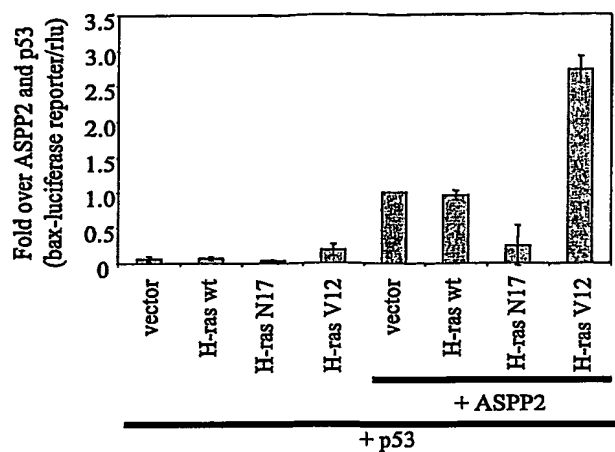


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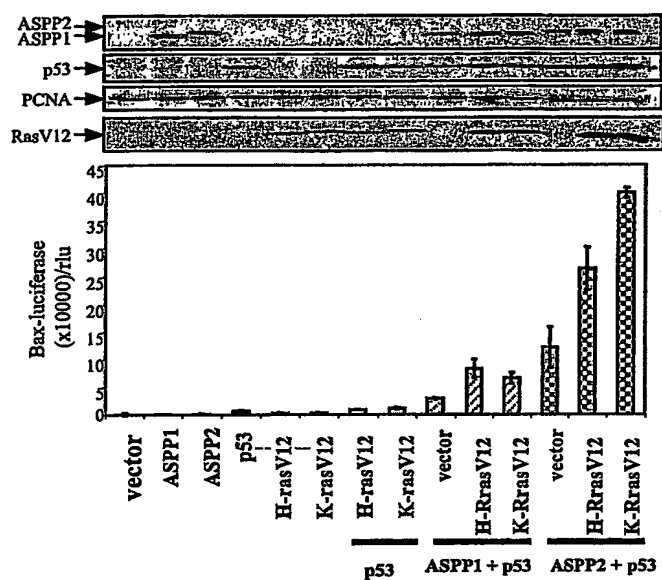
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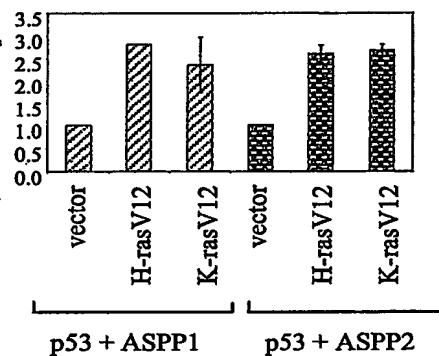


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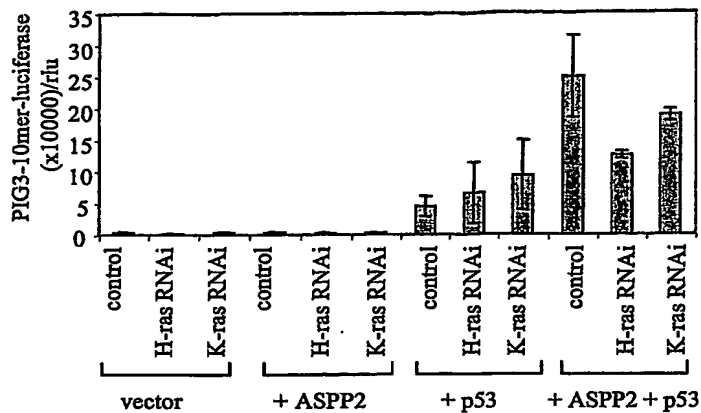
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fold over p53 + ASP1/2  
(bax-luciferase reporter)

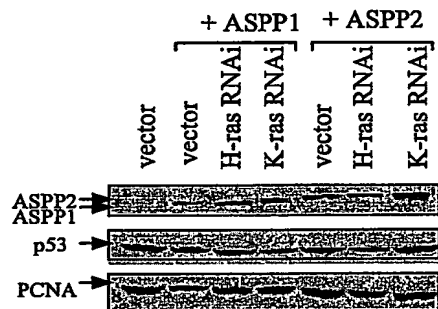
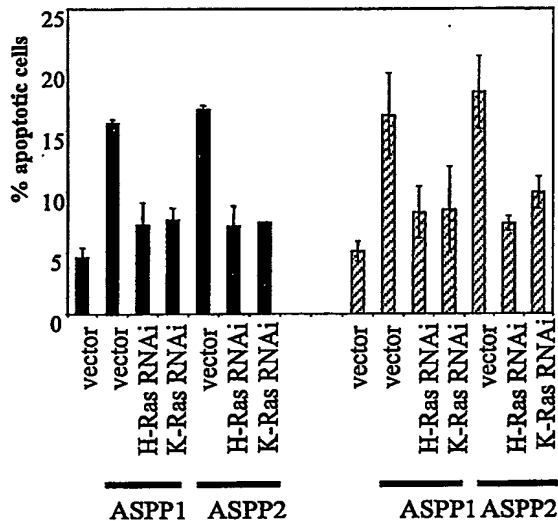


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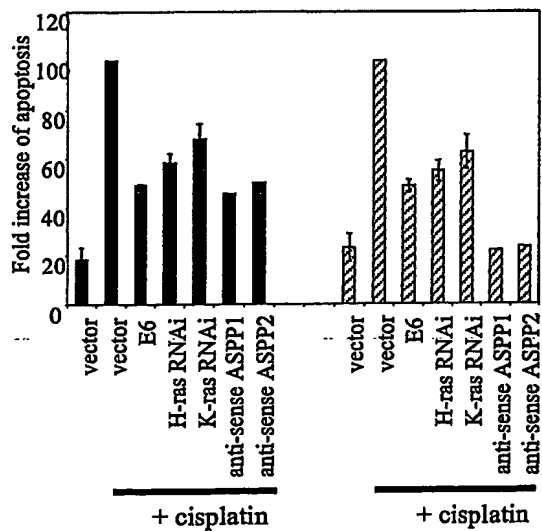
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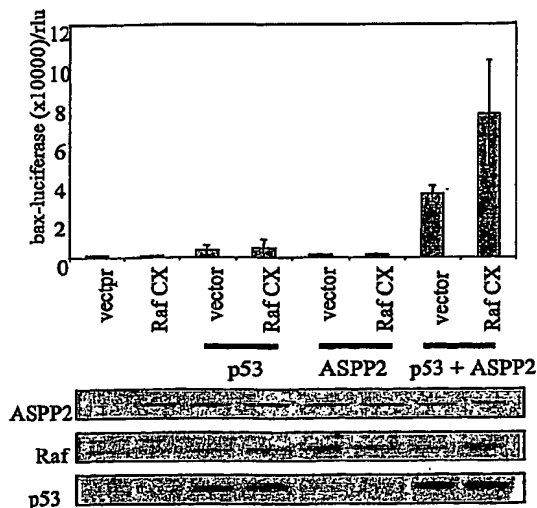


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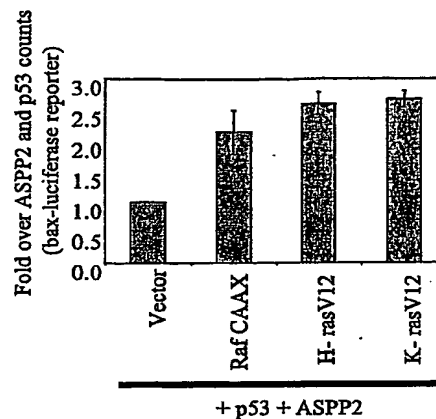
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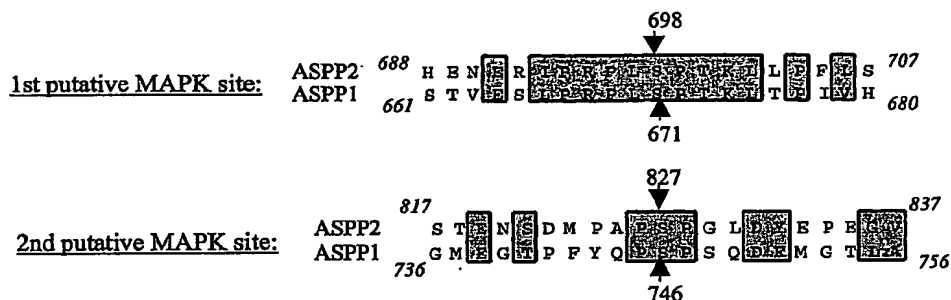
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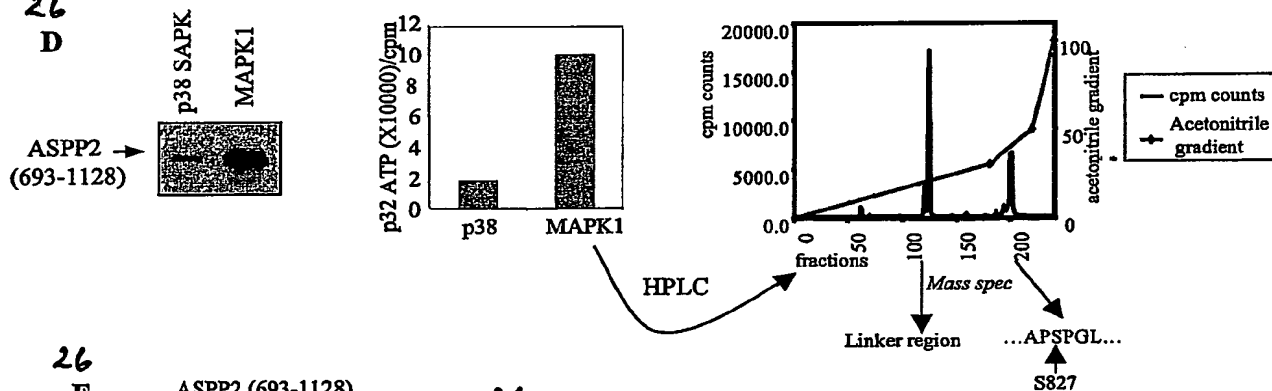
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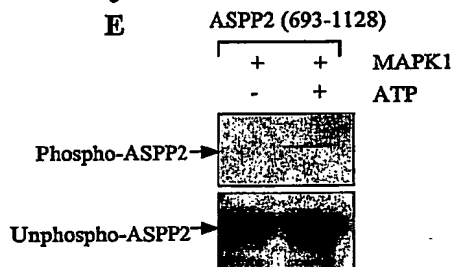
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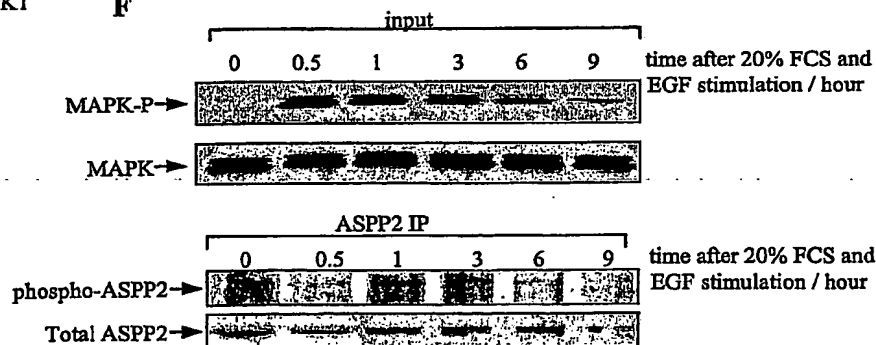
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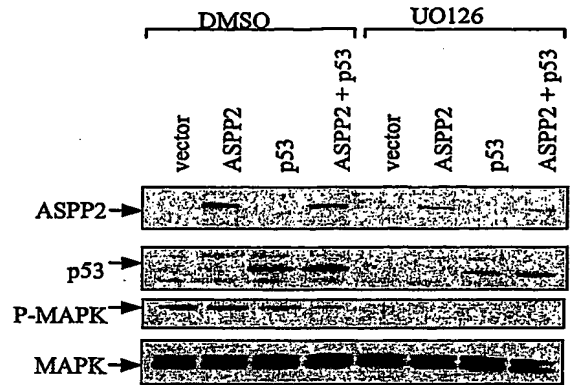
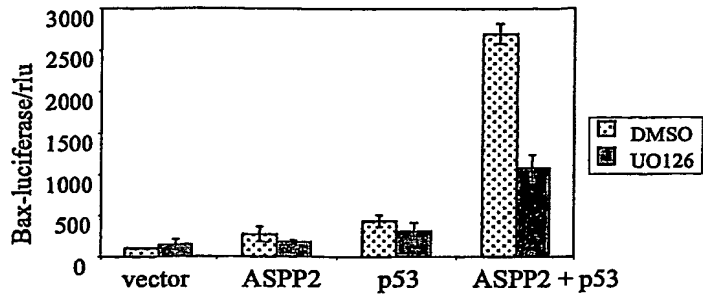
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E



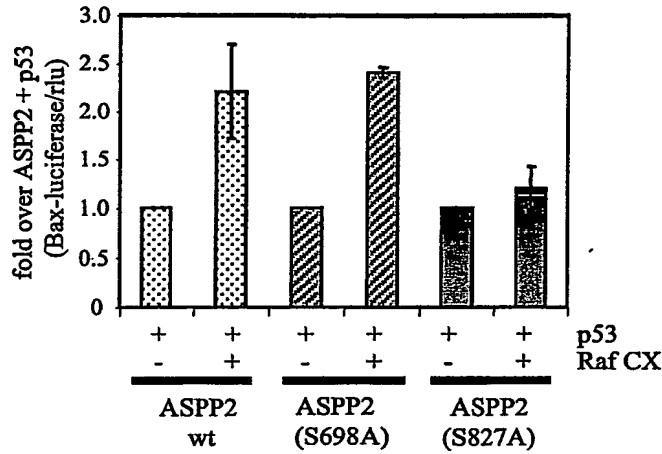
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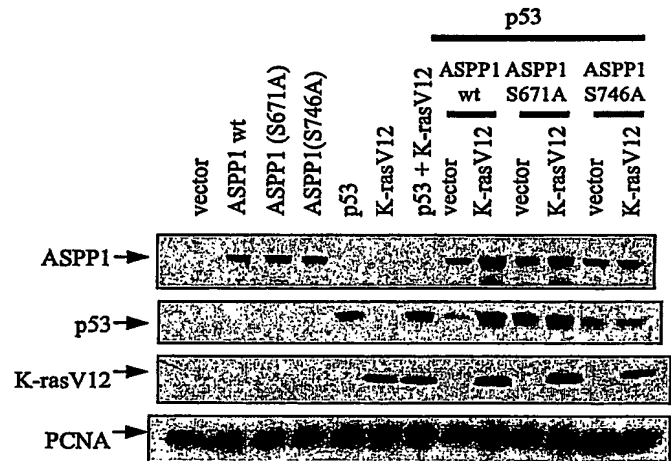
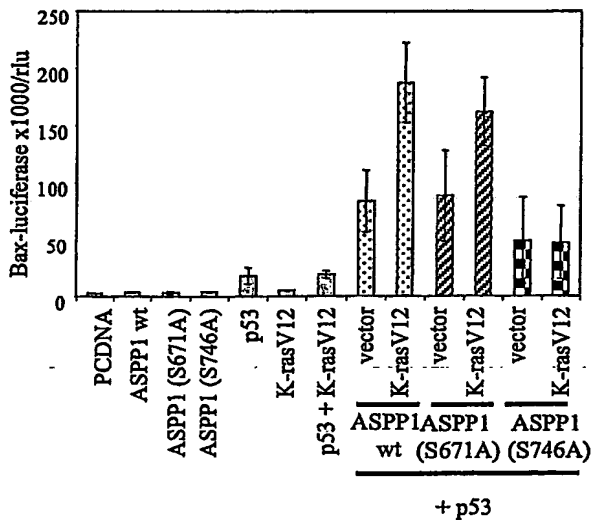
27  
A



27  
B

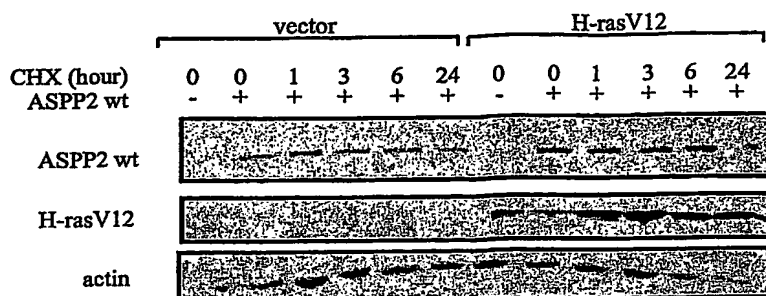


27  
C



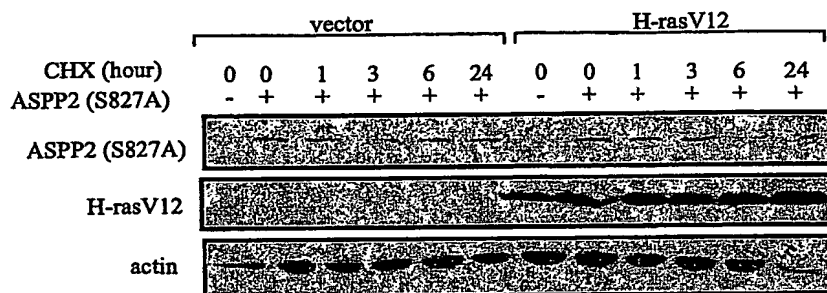
28

A



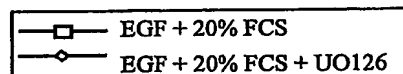
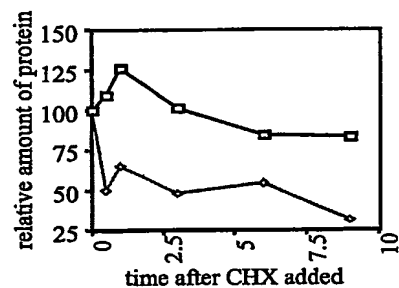
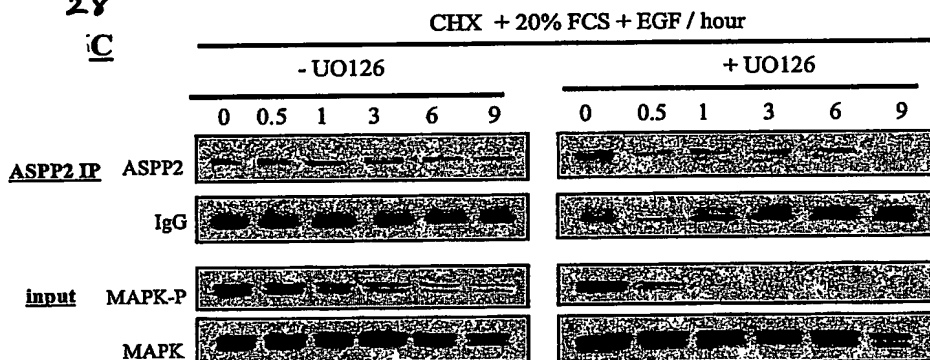
28

B



28

C



28

D

